

Figure 1a

1B3 VH chain, SEQ ID NO:1

QVQLKQSGAE LVRPGTSVKV SCKASGYAFT NYLIEWVKQR PGQGLEWIGV INPGSGGTNY
NEKFKDKATL TADKSSNTAY MQLSSLTSDD SAVYFCARRG WDYFDYWGQG TTLTVSS

Figure 1b

1B3 VL chain, SEQ ID NO:2

QIVLTQSPAL MSTSPGEKVT MTCSASSSVS YIYWFQKPG SSPKPWIYRT SNLASGVPAR
FSGSGSGTSY SLTISSMEAE DAATYYCQQW DNNPYTFGGG TKLEIK

Figure 2a

1B3(humanised) VH chain, SEQ ID NO:13

QVQLVQSGAE VKKPGASVKV SCKASGYAFT NYLIEWVRQA PGQGLEWIGV INPGSGGTNY
NEKFQGRVTL TADKSISTAY MELSRLRSDD TAVYYCARRG WDYFDYWGQG TLTVSS

Figure 2b

1B3(humanised) VL chain, SEQ ID NO:14

DIQLTQSPSS LSASVGDRVT ITCQASQDVS YIYWFQKPG KAPKPWIYRT SNLATGVPSR
FSGSGSGTDY TFTISLQPE DIATYYCQQW DNNPYTFGQG TKLEIK

Figure 3a

Mu 1	QVQLKQSGAELVRPGTSVKVSCASGYAFTNYLIEWVKRPGQGLEWIGV	50
	. :.. .	
Hu 1	QVQLVQSGAEVKKPGASVKVSCASGYAFTNYLIEWVRQAPGQGLEWIGV	50
51	INPGSGGTNYNEKFKDKATLTADKSSNTAYMQLSSLTSDDSAVYFCARRG	100
51	INPGSGGTNYNEKFQGRVTLTADKSISTAYMELSRLRSDTAVYYCARRG	100
101	WDYFDYWGQGTTLTVSS	117
101	WDYFDYWGQGTTLTVSS	117

Figure 3b

Mu 1	QIVLTQSPALMSTSPGEKVTMTCSASSSVSYIYWFQKPGSSPKPWIYRT	50
	. . :.. .	
Hu 1	DIQLTQSPSSLSASVGDRVTITCQASQDVSIIYWFQKPGKAPKPWIYRT	50
51	SNLASGVPARFSGSGGTSYSLTISSMEAEADAATYYCQQWDNNPYTFGGG	100
	:	
51	SNLATGVPSRFSGSGSGTDYFTFTISLQPEDATYYCQQWDNNPYTFGGG	100
101	TKLEIK	106
101	TKLEIK	106

Figure 4

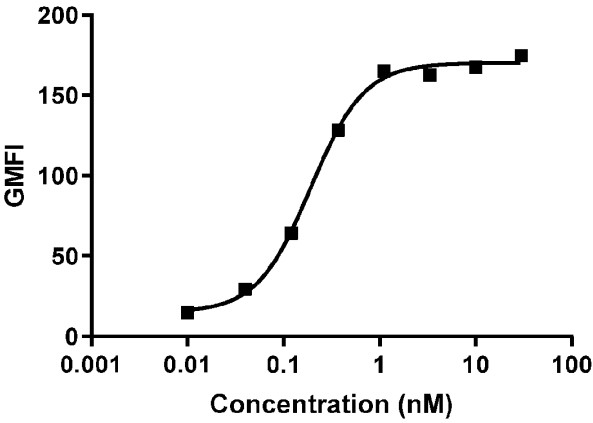


Figure 5

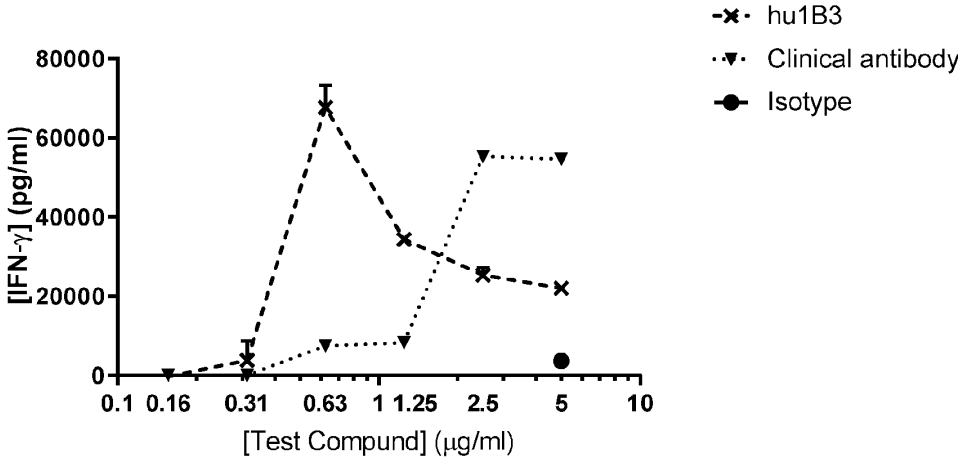


Figure 6

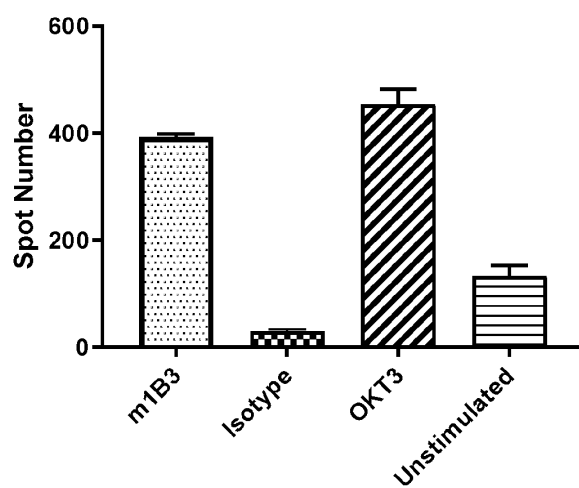


Figure 7

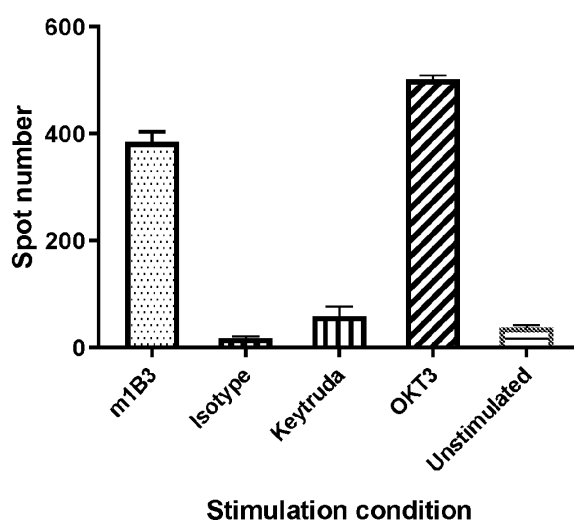


Figure 8

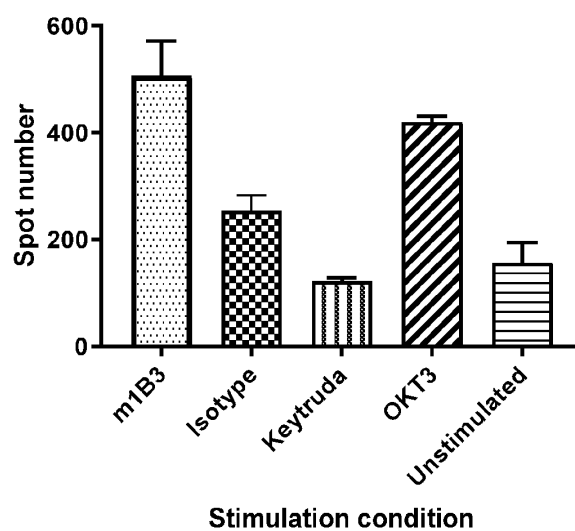


Figure 9

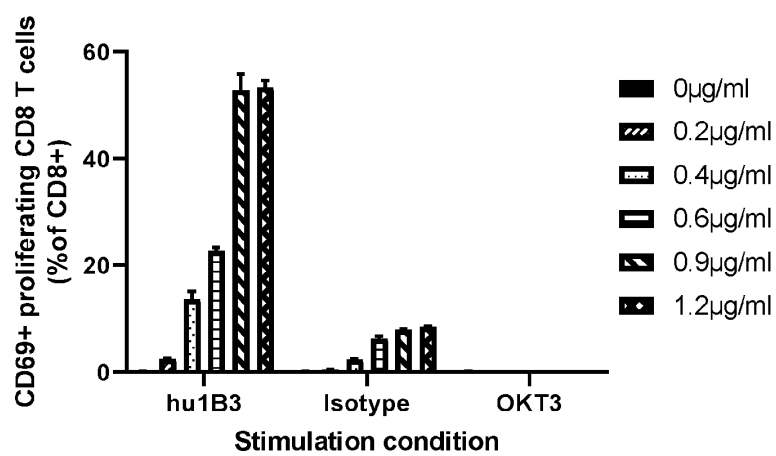


Figure 10

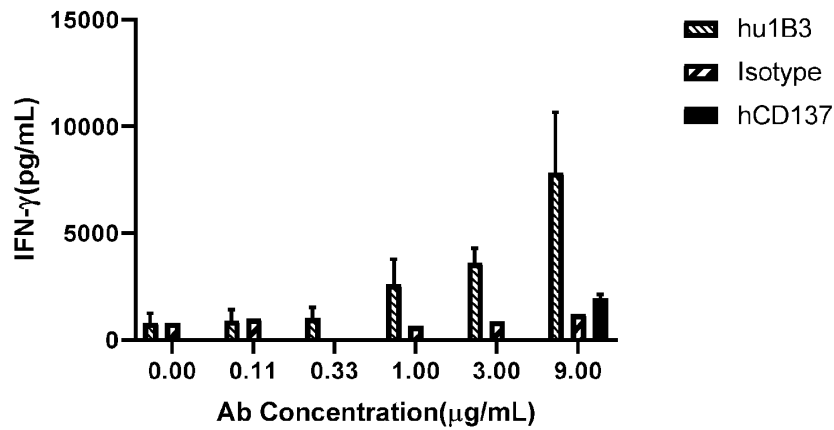


Figure 11

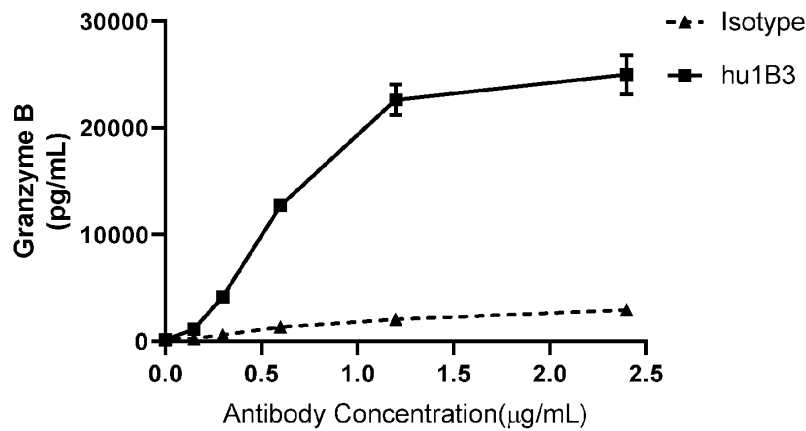


Figure 12

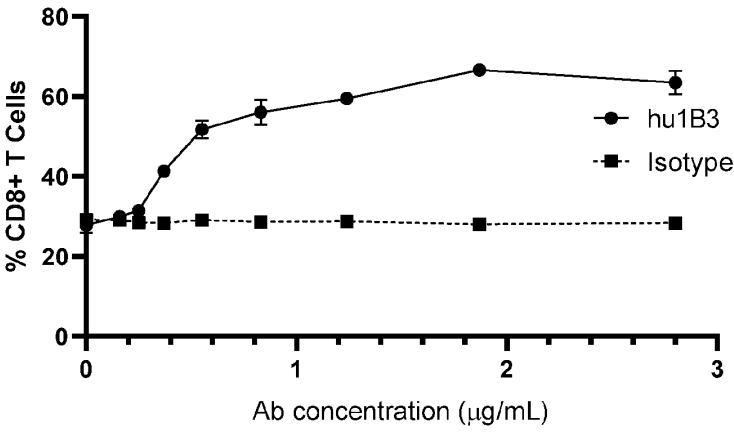


Figure 13

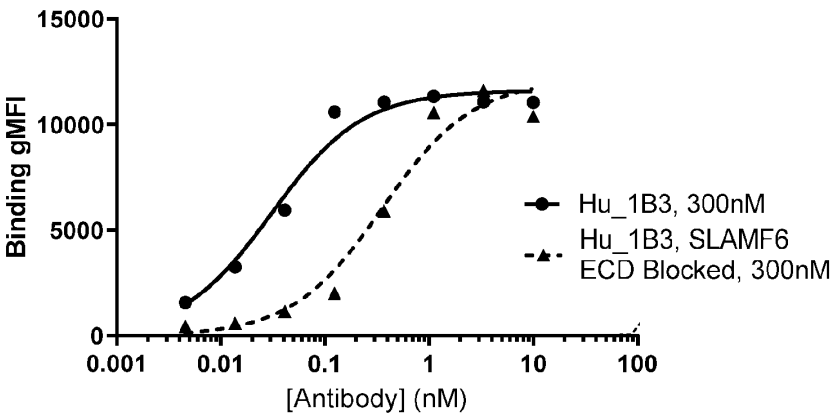


Figure 14

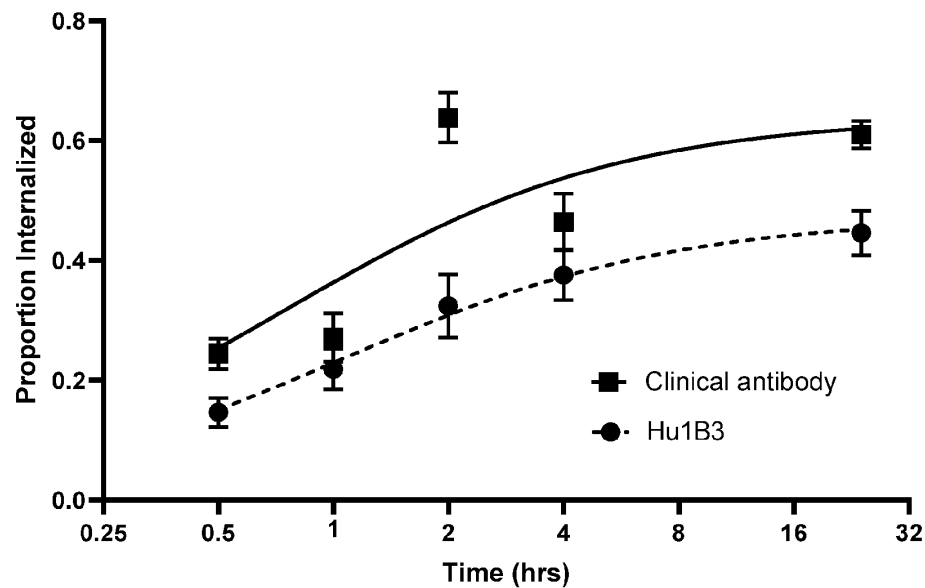


Figure 15

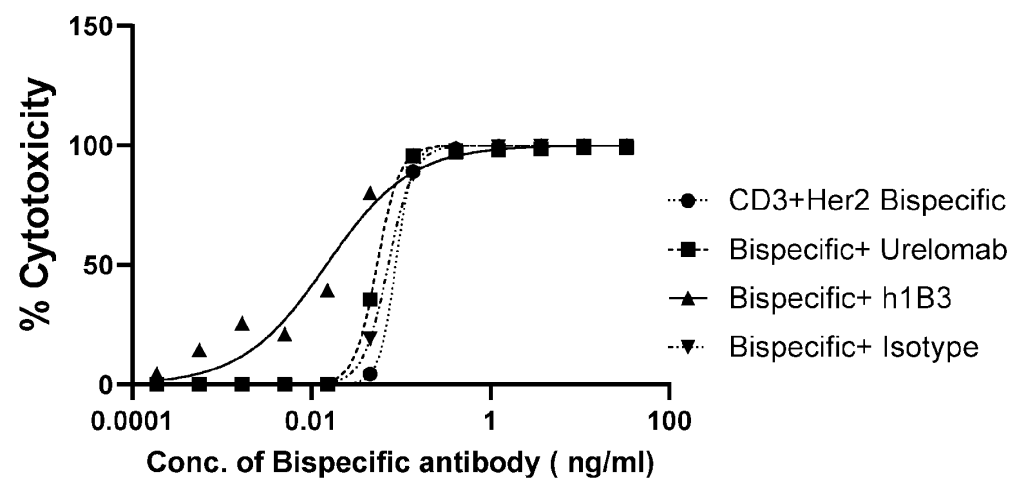


Figure 16

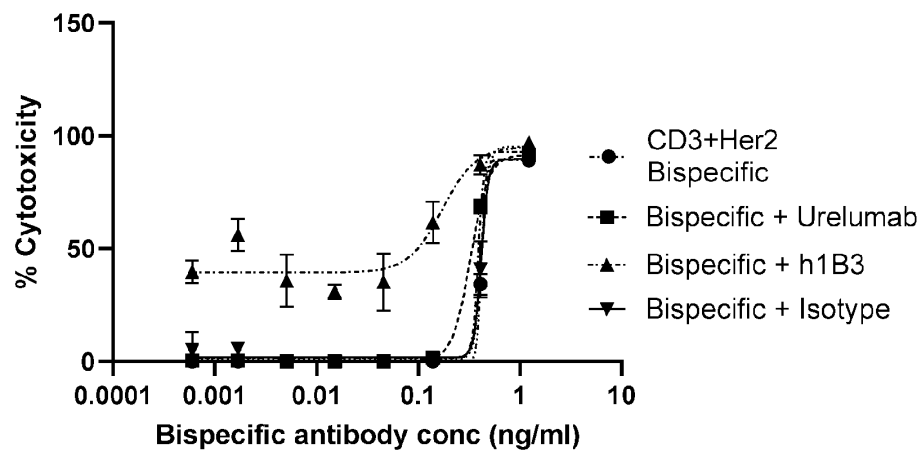


Figure17

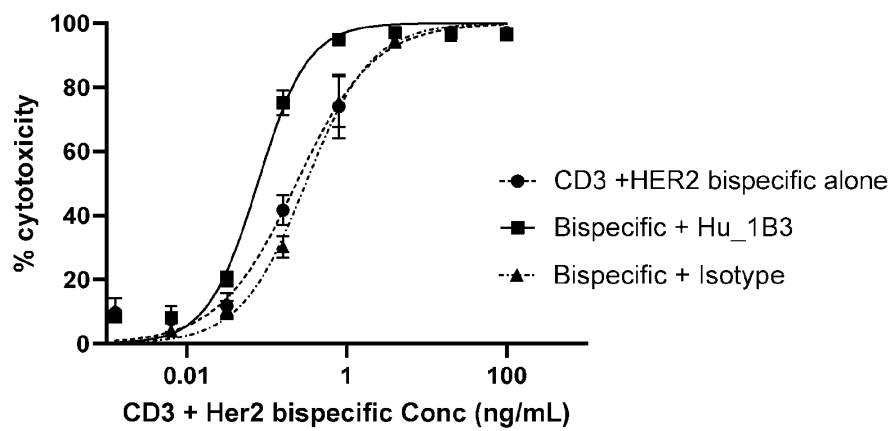


Figure 18

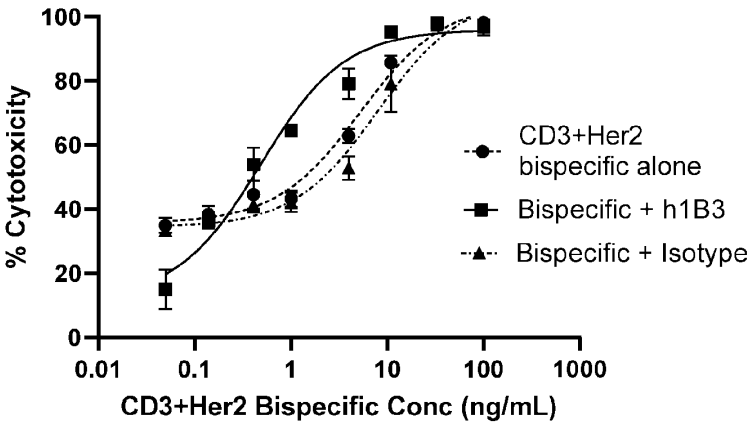


Figure 19

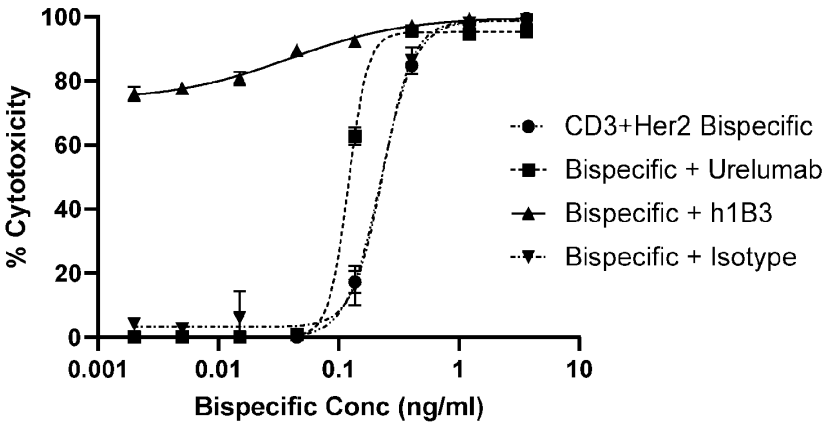


Figure 20

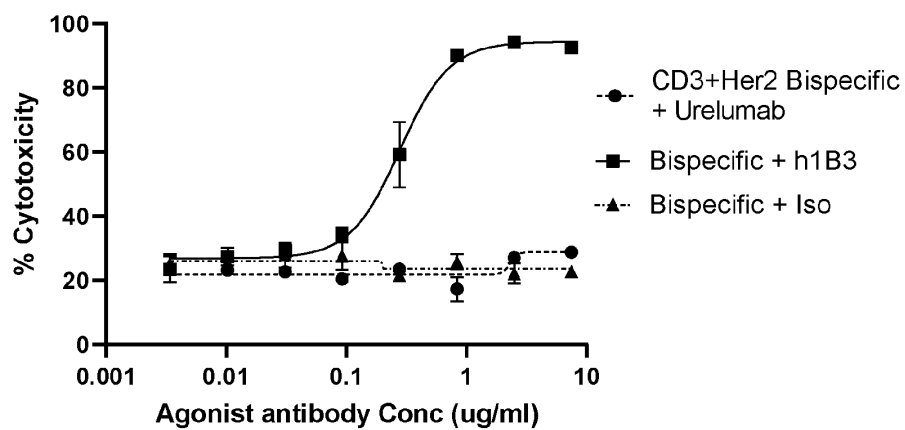


Figure 17a

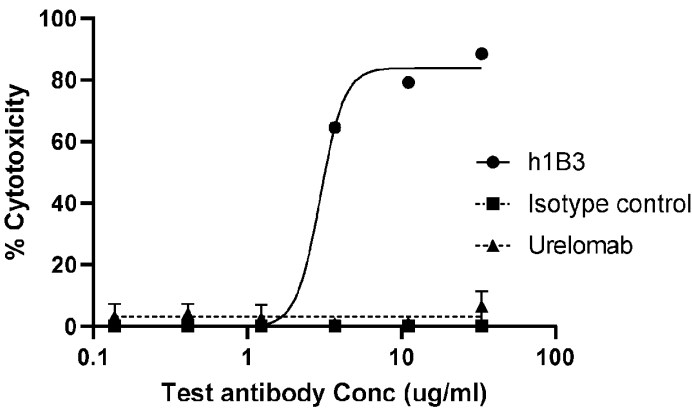


Figure 17b

